

Schwadron

CRF Errors Corrected by the STIC System, Branch

1644

Serial Number: 09/155,514

CRF Processing Date: 1/31/2003
Edited by: _____
Verified by: _____ (STIC staff)

Changed a file from non-ASCII to ASCII
 Changed the margins in cases where the sequence text was "wrapped" down to the next line.
 Edited a format error in the Current Application Data section, specifically:

ENTERED Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other _____.

Added the mandatory heading and subheadings for "Current Application Data".
 Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
 Changed the spelling of a mandatory field (the headings or subheadings), specifically:

Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:

Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: 1

Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

Inserted colons after headings/subheadings. Headings edited included:

Deleted extra, invalid, headings used by an applicant, specifically:

Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of file;
 page numbers throughout text; other invalid text, such as _____.

Inserted mandatory headings, specifically:

Corrected an obvious error in the response, specifically:

Edited identifiers where upper case is used but lower case is required, or vice versa.

Corrected an error in the Number of Sequences field, specifically:

A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

Other:

Input Set: I155514.RAW

This Raw Listing contains the General Information
Section and up to first 5 pages.

1 <110> APPLICANT: Kainoh, Mie
2 Tanaka, Toshiaki
3 <120> TITLE OF INVENTION: Chimeric proteins, their heterodimer complexes, and plate
4 substitutes
5 <130> FILE REFERENCE: 1102-98
6 <140> CURRENT APPLICATION NUMBER: US/09/155,514
7 <141> CURRENT FILING DATE: 1998-11-17
8 <150> EARLIER APPLICATION NUMBER: PCT/JP98/00370
9 <151> EARLIER FILING DATE: 1997-01-29
10 <150> EARLIER APPLICATION NUMBER: JP 9-15118
11 <151> EARLIER FILING DATE: 1997-01-29
12 <150> EARLIER APPLICATION NUMBER: JP 9-234544
13 <151> EARLIER FILING DATE: 1997-08-29
14 <160> NUMBER OF SEQ ID NOS: 31
15 <170> SOFTWARE: Wordperfect 8.0
16 <210> SEQ ID NO 1
17 <211> LENGTH: 4228
18 <212> TYPE: DNA
19 <213> ORGANISM: Homo sapien
20 <220> FEATURE:
21 <221> NAME/KEY: CDS
22 <222> LOCATION: 1...2958, 3316...3360, 3480...3808, 3905...4228
23 <300> PUBLICATION INFORMATION:
24 <400> SEQUENCE: 1
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-20 -15 -10
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Gly Val Pro Thr Gly Arg Pro Tyr Asn Val Asp Thr Glu Ser Ala Leu
-5 1 5
ctt tac cag ggc ccc cac aac acg ctg ttc ggc tac tcg gtc gtg ctg 192
Leu Tyr Gln Gly Pro His Asn Thr Leu Phe Gly Tyr Ser Val Val Leu
10 15 20 25
cac agc cac ggg gcg aac cga tgg ctc cta gtg ggt gcg ccc act gcc 240
His Ser His Gly Ala Asn Arg Trp Leu Leu Val Gly Ala Pro Thr Ala
30 35 40
aac tgg ctc gcc aac gct tca gtg atc aat ccc ggg gcg att tac aga 288
Asn Trp Leu Ala Asn Ala Ser Val Ile Asn Pro Gly Ala Ile Tyr Arg
45 50 55
tgc acg atc gga aag aat ccc ggc cag acg tgc gaa cag ctc cag ctg 336
Cys Arg Ile Gly Lys Asn Pro Gly Gln Thr Cys Glu Gln Leu Gln Leu
44

Input Set: I155514.RAW

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48	75	80	85	
49	gac aat cag tgg ttg ggg gtc aca ctt tcc aga cag cca gga gaa aat			432
50	Asp Asn Gln Trp Leu Gly Val Thr Leu Ser Arg Gln Pro Gly Glu Asn			
51	90	95	100	105
52	gga tcc atc gtg act tgt ggg cat aga tgg aaa aat ata ttt tac ata			480
53	Gly Ser Ile Val Thr Cys Gly His Arg Trp Lys Asn Ile Phe Tyr Ile			
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55	aag aat gaa aat aag ctc ccc act ggt ggt tgc tat gga gtg ccc cct			528
56	Lys Asn Glu Asn Lys Leu Pro Thr Gly Gly Cys Tyr Gly Val Pro Pro			
57	125	130	135	
58	gat tta cga aca gaa ctg agt aaa aga ata gct ccg tgc tat caa gat			576
59	Asp Leu Arg Thr Glu Leu Ser Lys Arg Ile Ala Pro Cys Tyr Gln Asp			
60	140	145	150	
61	tat gtg aaa aaa ttt gga gaa aat ttt gca tca tgc taa gct gga ata			624
62	Tyr Val Lys Lys Phe Gly Glu Asn Phe Ala Ser Cys Gln Ala Gly Ile			
63	155	160	165	
64	tcc agt ttt tac aca aag gat tta att gtg atg ggg gcc cca gga tca			672
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67	tct tac tgg act ggc tct ctt ttt gtc tac aat ata act aca aat aaa			720
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69	190	195	200	
70	tac aag gct ttt tta gac aaa caa aat caa gta aaa ttt gga agt tat			768
71	Tyr Lys Ala Phe Leu Asp Lys Gln Asn Gln Val Lys Phe Gly Ser Tyr			
72	205	210	215	
73	tta gga tat tca gtc gga gct ggt cat ttt cgg agc cag cat act acc			816
74	Leu Gly Tyr Ser Val Gly Ala Gly His Phe Arg Ser Gln His Thr Thr			
75	220	225	230	
76	gaa gta gtc gga gga gct cct caa cat gag cag att ggt aag gca tat			864
77	Glu Val Val Gly Gly Ala Pro Gln His Glu Gln Ile Gly Lys Ala Tyr			
78	235	240	245	
79	ata ttc agc att gat gaa aaa gaa cta aat atc tta cat gaa atg aaa			912
80	Ile Phe Ser Ile Asp Glu Lys Glu Leu Asn Ile Leu His Glu Met Lys			
81	250	255	260	265
82	ggt aaa aag ctt gga tcg tac ttt gga gct tct gtc tgc gac			960
83	Gly Lys Lys Leu Gly Ser Tyr Phe Gly Ala Ser Val Cys Ala Val Asp			
84	270	275	280	
85	ctc aat gca gat ggc ttc tca gat ctg ctc gtg gga gca ccc atg cag			1008
86	Leu Asn Ala Asp Gly Phe Ser Asp Leu Leu Val Gly Ala Pro Met Gln			
87	285	290	295	
88	agc acc atc aga gag gaa gga aga gtg ttt gtg tac atc aac tct ggc			1056
89	Ser Thr Ile Arg Glu Glu Gly Arg Val Phe Val Tyr Ile Asn Ser Gly			
90	300	305	310	
91	tcg gga gca gta atg aat gca atg gaa aca aac ctc gtt gga agt gac			1104
92	Ser Gly Ala Val Met Asn Ala Met Glu Thr Asn Leu Val Gly Ser Asp			
93	315	320	325	
94	aaa tat gct gca aga ttt ggg gaa tct ata gtt aat ctt ggc gac att			1152

Input Set: I155514.RAW

95	Lys	Tyr	Ala	Ala	Arg	Phe	Gly	Glu	Ser	Ile	Val	Asn	Leu	Gly	Asp	Ile	
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98	Asp	Asn	Asp	Gly	Phe	Glu	Asp	Val	Ala	Ile	Gly	Ala	Pro	Gln	Glu	Asp	
99					350					355					360		
100	gac	ttg	caa	ggt	gct	att	tat	att	tac	aat	ggc	cgt	gca	gat	ggg	atc	1248
101	Asp	Leu	Gln	Gly	Ala	Ile	Tyr	Ile	Tyr	Asn	Gly	Arg	Ala	Asp	Gly	Ile	
102					365				370			375					
103	tcg	tca	acc	ttc	tca	cag	aga	att	gaa	gga	ctt	cag	atc	agc	aaa	tcg	1296
104	Ser	Ser	Thr	Phe	Ser	Gln	Arg	Ile	Glu	Gly	Leu	Gln	Ile	Ser	Lys	Ser	
105					380				385			390					
106	tta	agt	atg	ttt	gga	cag	tct	ata	tca	gga	caa	att	gat	gca	gat	att	1344
107	Leu	Ser	Met	Phe	Gly	Gln	Ser	Ile	Ser	Gly	Gln	Ile	Asp	Ala	Asp	Asn	
108					395				400			405					
109	aat	ggc	tat	gta	gat	gta	gca	ggt	cgt	gct	ttt	cg	tct	gat	tct	gct	1392
110	Asn	Gly	Tyr	Val	Asp	Val	Ala	Val	Gly	Ala	Phe	Arg	Ser	Asp	Ser	Ala	
111					410				415			420			425		
112	gtc	ttg	cta	agg	aca	aga	cct	gta	gta	att	gtt	gac	gct	tct	tta	agc	1440
113	Val	Leu	Leu	Arg	Thr	Arg	Pro	Val	Val	Ile	Val	Asp	Ala	Ser	Leu	Ser	
114					430				435			440					
115	cac	cct	gag	tca	gta	aat	aga	acg	aaa	ttt	gac	tgt	gtt	gaa	aat	gga	1488
116	His	Pro	Glu	Ser	Val	Asn	Arg	Thr	Lys	Phe	Asp	Cys	Val	Glu	Asn	Gly	
117					445				450			455					
118	tgg	cct	tct	gtg	tgc	ata	gat	cta	aca	ctt	tgt	ttc	tca	tat	aag	ggc	1536
119	Trp	Pro	Ser	Val	Cys	Ile	Asp	Leu	Thr	Leu	Cys	Phe	Ser	Tyr	Lys	Gly	
120					460				465			470					
121	aag	gaa	gtt	cca	gg	tac	att	gtt	ttt	tat	aac	atg	agt	ttg	gat		1584
122	Lys	Glu	Val	Pro	Gly	Tyr	Ile	Val	Leu	Phe	Tyr	Asn	Met	Ser	Leu	Asp	
123					475				480			485					
124	gtg	aac	aga	aag	gca	gag	tct	cca	cca	aga	ttc	tat	ttc	tct	tct	aat	1632
125	Val	Asn	Arg	Lys	Ala	Glu	Ser	Pro	Pro	Arg	Phe	Tyr	Phe	Ser	Ser	Asn	
126					490				495			500			505		
127	gga	act	tct	gac	gtg	att	aca	gga	agc	ata	cag	gtg	tcc	agc	aga	gaa	1680
128	Gly	Thr	Ser	Asp	Val	Ile	Thr	Gly	Ser	Ile	Gln	Val	Ser	Ser	Arg	Glu	
129					510				515			520					
130	gct	aac	tgt	aga	aca	cat	caa	gca	ttt	atg	cg	aaa	gat	gtg	cg	gac	1728
131	Ala	Asn	Cys	Arg	Thr	His	Gln	Ala	Phe	Met	Arg	Lys	Asp	Val	Arg	Asp	
132					525				530			535					
133	atc	ctc	acc	cca	att	cag	att	gaa	gct	gct	tac	cac	ctt	ggt	cct	cat	1776
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135					540				545			550					
136	gtc	atc	agt	aaa	cga	agt	aca	gag	gaa	ttc	cca	cca	ctt	cag	cca	att	1824
137	Val	Ile	Ser	Lys	Arg	Ser	Thr	Glu	Glu	Phe	Pro	Pro	Leu	Gln	Pro	Ile	
138					555				560			565					
139	ctt	cag	cag	aag	aaa	gaa	aaa	gac	ata	atg	aaa	aaa	aca	ata	aac	ttt	1872
140	Leu	Gln	Gln	Lys	Lys	Glu	Lys	Asp	Ile	Met	Lys	Lys	Thr	Ile	Asn	Phe	
141					570				575			580			585		
142	gca	agg	ttt	tgt	gcc	cat	gaa	aat	tgt	tct	gct	gat	tta	cag	gtt	tct	1920
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Input Set: I155514.RAW

145	gca aag att ggg ttt ttg aag ccc cat gaa aat aaa aca tat ctt gct	1968
146	Ala Lys Ile Gly Phe Leu Lys Pro His Glu Asn Lys Thr Tyr Leu Ala	
147	605 610 615	
148	gtt ggg agt atg aag aca ttg atg ttg aat gtg tcc ttg ttt aat gct	2016
149	Val Gly Ser Met Lys Thr Leu Met Leu Asn Val Ser Leu Phe Asn Ala	
150	620 625 630	
151	gga gat gat gca tat gaa acg act cta cat gtc aaa cta ccc gtg ggt	2064
152	Gly Asp Asp Ala Tyr Glu Thr Thr Leu His Val Lys Leu Pro Val Gly	
153	635 640 645	
154	ctt tat ttc att aag att tta gag ctg gaa gag aag caa ata aac tgt	2112
155	Leu Tyr Phe Ile Lys Ile Leu Glu Leu Glu Lys Gln Ile Asn Cys	
156	650 655 660 665	
157	gaa gtc aca gat aac tct ggc gtg gta caa ctt gac tgc agt att ggc	2160
158	Glu Val Thr Asp Asn Ser Gly Val Val Gln Leu Asp Cys Ser Ile Gly	
159	670 675 680	
160	tat ata tat gta gat cat ctc tca agg ata gat att agc ttt ctc ctg	2208
161	Tyr Ile Tyr Val Asp His Leu Ser Arg Ile Asp Ile Ser Phe Leu Leu	
162	685 690 695	
163	gat gtg agc tca ctc agc aga gcg gaa gag gac ctc agt atc aca gtg	2256
164	Asp Val Ser Ser Leu Ser Arg Ala Glu Glu Asp Leu Ser Ile Thr Val	
165	700 705 710	
166	cat gct acc tgt gaa aat gaa gag gaa atg gac aat cta aag cac agc	2304
167	His Ala Thr Cys Glu Asn Glu Glu Met Asp Asn Leu Lys His Ser	
168	715 720 725	
169	aga gtg act gta gca ata cct tta aaa tat gag gtt aag ctg act gtt	2352
170	Arg Val Thr Val Ala Ile Pro Leu Lys Tyr Glu Val Lys Leu Thr Val	
171	730 735 740 745	
172	cat ggg ttt gta aac cca act tca ttt gtg tat gga tca aat gat gaa	2400
173	His Gly Phe Val Asn Pro Thr Ser Phe Val Tyr Gly Ser Asn Asp Glu	
174) 750 755 760	
175	aat gag cct gaa acg tgc atg gtg gag aaa atg aac tta act ttc cat	2448
176	Asn Glu Pro Glu Thr Cys Met Val Glu Lys Met Asn Leu Thr Phe His	
177	765 770 775	
178	gtt atc aac act ggc aat agt atg gct ccc aat gtt agt gtg gaa ata	2496
179	Val Ile Asn Thr Gly Asn Ser Met Ala Pro Asn Val Ser Val Glu Ile	
180	780 785 790	
181	atg gta cca aat tct ttt agc ccc caa act gat aag ctg ttc aac att	2544
182	Met Val Pro Asn Ser Phe Ser Pro Gln Thr Asp Lys Leu Phe Asn Ile	
183	795 800 805	
184	ttg gat gtc cag act act act gga gaa tgc cac ttt gaa aat tat caa	2592
185	Leu Asp Val Gln Thr Thr Gly Glu Cys His Phe Glu Asn Tyr Gln	
186	810 815 820 825	
187	aga gtg tgt gca tta gag cag caa aag agt gca atg cag acc ttg aaa	2640
188	Arg Val Cys Ala Leu Glu Gln Gln Lys Ser Ala Met Gln Thr Leu Lys	
189	830 835 840	
190	ggc ata gtc cgg ttc ttg tcc aag act gat aag agg cta ttg tac tgc	2688
191	Gly Ile Val Arg Phe Leu Ser Lys Thr Asp Lys Arg Leu Leu Tyr Cys	
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193	ata aaa gct gat cca cat tgt tta aat ttc ttg tgt aat ttt ggg aaa	2736
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Input Set: I155514.RAW

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198	875	880	885	
199	cca tcc att tta gaa atg gat gag act tca gca ctc aag ttt gaa ata			2832
200	Pro Ser Ile Leu Glu Met Asp Glu Thr Ser Ala Leu Lys Phe Glu Ile			
201	890	895	900	905
202	aga gca aca ggt ttt cca gag cca aat cca aga gta att gaa cta aac			2880
203	Arg Ala Thr Gly Phe Pro Glu Pro Asn Pro Arg Val Ile Glu Leu Asn			
204	910	915	920	
205	aag gat gag aat gtt gcg cat gtt cta ctg gaa gga cta cat cat caa			2928
206	Lys Asp Glu Asn Val Ala His Val Leu Leu Glu Gly Leu His His Gln			
207	925	930	935	
208	aga ccc aaa cgt tat ttc acg gat ccc gag ctgctggaaag caggctcagc			2978
209	Arg Pro Lys Arg Tyr Phe Thr Asp Pro Glu			
210	940	945		
211	gctcctgcct ggacgcattcc cggctatgca gccccagtcc agggcagcaa ggcaggcccc	3038		
212	gtctgcctct tcacccggag cctctgcccc ccccaactcat gtcaggag agggtcttct	3098		
213	ggcttttcc caggctctgg gcaggcacag gctaggtgcc cctaaccag gccctgcaca	3158		
214	caaaggggca ggtgctgggc tcagacctgc caagagccat atccgggagg accctgcccc	3218		
215	tgacctaagg ccacccaaa ggccaaactc tccactccct cagctcggac accttctctc	3278		
216	ctcccgatt ccagtaactc ccaatcttct ctctgca gag ccc aaa tct tgt gac	3333		
217	Glu Pro Lys Ser Cys Asp			
218	950			
219	aaa act cac aca tgc cca ccg tgc cca ggtaagccag cccaggcctc	3380		
220	Lys Thr His Thr Cys Pro Pro Cys Pro			
221	955	960		
222	gccctccagc tcaaggcggg acaggtgccc tagagtaggcc tgcatccagg gacaggcccc	3440		
223	agccgggtgc tgacacgtcc acctccatct cttcctca gca cct gaa ctc ctg	3493		
224	Ala Pro Glu Leu Leu			
225	965			
226	ggg gga ccg tca gtc ttc ctc ttc cca aaa ccc aag gac acc ctc	3541		
227	Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu			
228	970	975	980	
229	atg atc tcc cgg acc cct gag gtc aca tgc gtg gtg gac gtg agc	3589		
230	Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser			
231	985	990	995	
232	cac gaa gac cct gag gtc aag ttc aac tgg tac gtg gac ggc gtg gag	3637		
233	His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu			
234	1000	1005	1010	1015
235	gtg cat aat gcc aag aca aag ccg cgg gag gag cag tac aac agc acg	3685		
236	Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr			
237	1020	1025	1030	
238	tac cgg gtg gtc agc gtc ctc acc gtc ctg cac cag gac tgg ctg aat	3733		
239	Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn			
240	1035	1040	1045	
241	ggc aag gag tac aag tgc aag gtc tcc aac aaa gcc ctc cca gcc ccc	3781		
242	Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro			
243	1050	1055	1060	
244	atc gag aaa acc atc tcc aaa gcc aaa ggtgggaccc gtggggtgcg	3828		

PAGE: 6

VERIFICATION SUMMARY
PATENT APPLICATION US/09/155,514

DATE: 02/03/2000

TIME: 13:21:00

Input Set: I155514.RAW

Line ? Error/Warning

Original Text

Input Set: I155514.RAW

PREVIOUSLY ERRORED SEQUENCES - EDITED

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12 -35 -30 -25
13 ggc ccc gaa gct gca ctc cgg gag acg gtg atg ctg ttg ctg tgc ctg 96
14 Gly Pro Glu Ala Ala Leu Arg Glu Thr Val Met Leu Leu Leu Cys Leu
15 -20 -15 -10
16 ggg gtc ccg acc ggc agg cct tac aac gtg gac act gag agc gcg ctg 144
17 Gly Val Pro Thr Gly Arg Pro Tyr Asn Val Asp Thr Glu Ser Ala Leu
18 -5 1 5
19 ctt tac cag ggc ccc cac aac acg ctg ttc ggc tac tcg gtc gtg ctg 192
20 Leu Tyr Gln Gly Pro His Asn Thr Leu Phe Gly Tyr Ser Val Val Leu
21 10 15 20 25
22 cac agc cac ggg gcg aac cga tgg ctc cta gtg ggt gcg ccc act gcc 240
23 His Ser His Gly Ala Asn Arg Trp Leu Leu Val Gly Ala Pro Thr Ala
24 30 35 40
25 aac tgg ctc gcc aac gct tca gtg atc aat ccc ggg gcg att tac aga 288
26 Asn Trp Leu Ala Asn Ala Ser Val Ile Asn Pro Gly Ala Ile Tyr Arg
27 45 50 55
28 tgc acg atc gga aag aat ccc ggc cag acg tgc gaa cag ctc cag ctg 336
29 Cys Arg Ile Gly Lys Asn Pro Gly Gln Thr Cys Glu Gln Leu Gln Leu
30 60 65 70
31 ggt agc cct aat gga gaa cct tgt gga aag act tgt ttg gaa gag aga 384
32 Gly Ser Pro Asn Gly Glu Pro Cys Gly Lys Thr Cys Leu Glu Glu Arg
33 75 80 85
34 gac aat cag tgg ttg ggg gtc aca ctt tcc aga cag cca gga gaa aat 432
35 Asp Asn Gln Trp Leu Gly Val Thr Leu Ser Arg Gln Pro Gly Glu Asn
36 90 95 100 105
37 gga tcc atc gtg act tgt ggg cat aga tgg aaa aat ata ttt tac ata 480
38 Gly Ser Ile Val Thr Cys Gly His Arg Trp Lys Asn Ile Phe Tyr Ile
39 110 115 120
40 aag aat gaa aat aag ctc ccc act ggt ggt tgc tat gga gtg ccc cct 528
41 Lys Asn Glu Asn Lys Leu Pro Thr Gly Gly Cys Tyr Gly Val Pro Pro
42 125 130 135
43 gat tta cga aca gaa ctg agt aaa aga ata gct ccg tgt tat caa gat 576
44 Asp Leu Arg Thr Glu Leu Ser Lys Arg Ile Ala Pro Cys Tyr Gln Asp
45 140 145 150
46 tat gtg aaa aaa ttt gga gaa aat ttt gca tca tgt caa gct gga ata 624
47 Tyr Val Lys Lys Phe Gly Glu Asn Phe Ala Ser Cys Gln Ala Gly Ile

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48

155

160

165

49

tcc agt ttt tac aca aag gat tta att gtg atg ggg gcc cca gga tca
Ser Ser Phe Tyr Thr Lys Asp Leu Ile Val Met Gly Ala Pro Gly Ser

672

50

Input Set: I155514.RAW

51	170	175	180	185	
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54	190	195	200		
55	tac aag gct ttt tta gac aaa caa aat caa gta aaa ttt gga agt tat				768
56	Tyr Lys Ala Phe Leu Asp Lys Gln Asn Gln Val Lys Phe Gly Ser Tyr				
57	205	210	215		
58	tta gga tat tca gtc gga gct ggt cat ttt cgg agc cag cat act acc				816
59	Leu Gly Tyr Ser Val Gly Ala Gly His Phe Arg Ser Gln His Thr Thr				
60	220	225	230		
61	gaa gta gtc gga gga gct cct caa cat gag cag att ggt aag gca tat				864
62	Glu Val Val Gly Gly Ala Pro Gln His Glu Gln Ile Gly Lys Ala Tyr				
63	235	240	245		
64	ata ttc agc att gat gaa aaa gaa cta aat atc tta cat gaa atg aaa				912
65	Ile Phe Ser Ile Asp Glu Lys Glu Leu Asn Ile Leu His Glu Met Lys				
66	250	255	260	265	
67	ggt aaa aag ctt gga tcg tac ttt gga gct tct gtc tgt gct gtg gac				960
68	Gly Lys Lys Leu Gly Ser Tyr Phe Gly Ala Ser Val Cys Ala Val Asp				
69	270	275	280		
70	ctc aat gca gat ggc ttc tca gat ctg ctc gtg gga gca ccc atg cag				1008
71	Leu Asn Ala Asp Gly Phe Ser Asp Leu Leu Val Gly Ala Pro Met Gln				
72	285	290	295		
73	agc acc atc aga gag gaa gga aga gtg ttt gtg tac atc aac tct ggc				1056
74	Ser Thr Ile Arg Glu Glu Gly Arg Val Phe Val Tyr Ile Asn Ser Gly				
75	300	305	310		
76	tcg gga gca gta atg aat gca atg gaa aca aac ctc gtt gga agt gac				1104
77	Ser Gly Ala Val Met Asn Ala Met Glu Thr Asn Leu Val Gly Ser Asp				
78	315	320	325		
79	aaa tat gct gca aga ttt ggg gaa tct ata gtt aat ctt ggc gac att				1152
80	Lys Tyr Ala Ala Arg Phe Gly Glu Ser Ile Val Asn Leu Gly Asp Ile				
81	330	335	340	345	
82	gac aat gat ggc ttt gaa gat gtt gct atc gga gct cca caa gaa gat				1200
83	Asp Asn Asp Gly Phe Glu Asp Val Ala Ile Gly Ala Pro Gln Glu Asp				
84	350	355	360		
85	gac ttg caa ggt gct att tat att tac aat ggc cgt gca gat ggg atc				1248
86	Asp Leu Gln Gly Ala Ile Tyr Ile Tyr Asn Gly Arg Ala Asp Gly Ile				
87	365	370	375		
88	tcg tca acc ttc tca cag aga att gaa gga ctt cag atc agc aaa tcg				1296
89	Ser Ser Thr Phe Ser Gln Arg Ile Glu Gly Leu Gln Ile Ser Lys Ser				
90	380	385	390		
91	tta agt atg ttt gga cag tac tct ata tca gga caa att gat gca gat att				1344
92	Leu Ser Met Phe Gly Gln Ser Ile Ser Gly Gln Ile Asp Ala Asp Asn				
93	395	400	405		
94	aat ggc tat gta gat gta gca ggt cgt gct ttt cgg tct gat tct gct				1392
95	Asn Gly Tyr Val Asp Val Ala Val Gly Ala Phe Arg Ser Asp Ser Ala				
96	410	415	420	425	
97	gtc ttg cta agg aca aga cct gta gta att gtt gac gct tct tta agc				1440
98	Val Leu Leu Arg Thr Arg Pro Val Val Ile Val Asp Ala Ser Leu Ser				
99	430	435	440		
100	cac cct gag tca gta aat aga acg aaa ttt gac tgt gtt gaa aat gga				1488

Input Set: I155514.RAW

101	His Pro Glu Ser Val Asn Arg Thr Lys Phe Asp Cys Val Glu Asn Gly			
102	445	450	455	
103	tgg cct tct gtg tgc ata gat cta aca ctt tgt ttc tca tat aag ggc		1536	
104	Trp Pro Ser Val Cys Ile Asp Leu Thr Leu Cys Phe Ser Tyr Lys Gly			
105	460	465	470	
106	aag gaa gtt cca ggt tac att gtt ttg ttt tat aac atg agt ttg gat		1584	
107	Lys Glu Val Pro Gly Tyr Ile Val Leu Phe Tyr Asn Met Ser Leu Asp			
108	475	480	485	
109	gtg aac aga aag gca gag tct cca cca aga ttc tat ttc tct tct aat		1632	
110	Val Asn Arg Lys Ala Glu Ser Pro Pro Arg Phe Tyr Phe Ser Ser Asn			
111	490	495	500	505
112	gga act tct gac gtg att aca gga agc ata cag gtg tcc agc aga gaa		1680	
113	Gly Thr Ser Asp Val Ile Thr Gly Ser Ile Gln Val Ser Ser Arg Glu			
114	510	515	520	
115	gct aac tgt aga aca cat caa gca ttt atg cgg aaa gat gtg cgg gac		1728	
116	Ala Asn Cys Arg Thr His Gln Ala Phe Met Arg Lys Asp Val Arg Asp			
117	525	530	535	
118	atc ctc acc cca att cag att gaa gct tac cac ctt ggt cct cat		1776	
119	Ile Leu Thr Pro Ile Gln Ile Glu Ala Ala Tyr His Leu Gly Pro His			
120	540	545	550	
121	gtc atc agt aaa cga agt aca gag gaa ttc cca cca ctt cag cca att		1824	
122	Val Ile Ser Lys Arg Ser Thr Glu Glu Phe Pro Pro Leu Gln Pro Ile			
123	555	560	565	
124	ctt cag cag aag aaa gaa aaa gac ata atg aaa aaa aca ata aac ttt		1872	
125	Leu Gln Gln Lys Lys Glu Lys Asp Ile Met Lys Lys Thr Ile Asn Phe			
126	570	575	580	585
127	gca agg ttt tgt gcc cat gaa aat tgt tct gct gat tta cag gtt tct		1920	
128	Ala Arg Phe Cys Ala His Glu Asn Cys Ser Ala Asp Leu Gln Val Ser			
129	590	595	600	
130	gca aag att ggg ttt ttg aag ccc cat gaa aat aaa aca tat ctt gct		1968	
131	Ala Lys Ile Gly Phe Leu Lys Pro His Glu Asn Lys Thr Tyr Leu Ala			
132	605	610	615	
133	gtt ggg agt atg aag aca ttg atg ttg aat gtg tcc ttg ttt aat gct		2016	
134	Val Gly Ser Met Lys Thr Leu Met Leu Asn Val Ser Leu Phe Asn Ala			
135	620	625	630	
136	gga gat gat gca tat gaa acg act cta cat gtc aaa cta ccc gtg ggt		2064	
137	Gly Asp Asp Ala Tyr Glu Thr Thr Leu His Val Lys Leu Pro Val Gly			
138	635	640	645	
139	ctt tat ttc att aag att tta gag ctg gaa gag aag caa ata aac tgt		2112	
140	Leu Tyr Phe Ile Lys Ile Leu Glu Leu Glu Glu Lys Gln Ile Asn Cys			
141	650	655	660	665
142	gaa gtc aca gat aac tct ggc gtg gta caa ctt gac tgc agt att ggc		2160	
143	Glu Val Thr Asp Asn Ser Gly Val Val Gln Leu Asp Cys Ser Ile Gly			
144	670	675	680	
145	tat ata tat gta gat cat ctc tca agg ata gat att agc ttt ctc ctg		2208	
146	Tyr Ile Tyr Val Asp His Leu Ser Arg Ile Asp Ile Ser Phe Leu Leu			
147	685	690	695	
148	gat gtg agc tca ctc agc aga gcg gaa gag gac ctc agt atc aca gtg		2256	
149	Asp Val Ser Ser Leu Ser Arg Ala Glu Glu Asp Leu Ser Ile Thr Val			
150	700	705	710	

Input Set: I155514.RAW

151	cat gct acc tgt gaa aat gaa gag gaa atg gac aat cta aag cac agc	2304
152	His Ala Thr Cys Glu Asn Glu Glu Glu Met Asp Asn Leu Lys His Ser	
153	715 720 725	
154	aga gtg act gta gca ata cct tta aaa tat gag gtt aag ctg act gtt	2352
155	Arg Val Thr Val Ala Ile Pro Leu Lys Tyr Glu Val Lys Leu Thr Val	
156	730 735 740 745	
157	cat ggg ttt gta aac cca act tca ttt gtg tat gga tca aat gat gaa	2400
158	His Gly Phe Val Asn Pro Thr Ser Phe Val Tyr Gly Ser Asn Asp Glu	
159	750 755 760	
160	aat gag cct gaa acg tgc atg gtg gag aaa atg aac tta act ttc cat	2448
161	Asn Glu Pro Glu Thr Cys Met Val Glu Lys Met Asn Leu Thr Phe His	
162	765 770 775	
163	gtt atc aac act ggc aat agt atg gct ccc aat gtt agt gtg gaa ata	2496
164	Val Ile Asn Thr Gly Asn Ser Met Ala Pro Asn Val Ser Val Glu Ile	
165	780 785 790	
166	atg gta cca aat tct ttt agc ccc caa act gat aag ctg ttc aac att	2544
167	Met Val Pro Asn Ser Phe Ser Pro Gln Thr Asp Lys Leu Phe Asn Ile	
168	795 800 805	
169	ttg gat gtc cag act act act gga gaa tgc cac ttt gaa aat tat caa	2592
170	Leu Asp Val Gln Thr Thr Gly Glu Cys His Phe Glu Asn Tyr Gln	
171	810 815 820 825	
172	aga gtg tgt gca tta gag cag caa aag agt gca atg cag acc ttg aaa	2640
173	Arg Val Cys Ala Leu Glu Gln Gln Lys Ser Ala Met Gln Thr Leu Lys	
174	830 835 840	
175	ggc ata gtc cgg ttc ttg tcc aag act gat aag agg cta ttg tac tgc	2688
176	Gly Ile Val Arg Phe Leu Ser Lys Thr Asp Lys Arg Leu Leu Tyr Cys	
177	845 850 855	
178	ata aaa gct gat cca cat tgt tta aat ttc ttg tgt aat ttt ggg aaa	2736
179	Ile Lys Ala Asp Pro His Cys Leu Asn Phe Leu Cys Asn Phe Gly Lys	
180	860 865 870	
181	atg gaa agt gga aaa gaa gcc agt gtt cat atc caa ctg gaa ggc cgg	2784
182	Met Glu Ser Gly Lys Glu Ala Ser Val His Ile Gln Leu Glu Gly Arg	
183	875 880 885	
184	cca tcc att tta gaa atg gat gag act tca gca ctc aag ttt gaa ata	2832
185	Pro Ser Ile Leu Glu Met Asp Glu Thr Ser Ala Leu Lys Phe Glu Ile	
186	890 895 900 905	
187	aga gca aca ggt ttt cca gag cca aat cca aga gta att gaa cta aac	2880
188	Arg Ala Thr Gly Phe Pro Glu Pro Asn Pro Arg Val Ile Glu Leu Asn	
189	910 915 920	
190	aag gat gag aat gtt gcg cat gtt cta ctg gaa gga cta cat cat caa	2928
191	Lys Asp Glu Asn Val Ala His Val Leu Leu Glu Gly Leu His His Gln	
192	925 930 935	
193	aga ccc aaa cgt tat ttc acg gat ccc gag ctgctgaaag caggctcagc	2978
194	Arg Pro Lys Arg Tyr Phe Thr Asp Pro Glu	
195	940 945	
196	gctcctgcct ggacgcatcc cggctatgca gccccagtcc agggcagcaa ggcaggcccc	3038
197	gtctgcctct tcacccggag cctctgccc cccactcat gtcaggag agggcttct	3098
198	ggcttttcc caggctctgg gcaggcacag gctagggtgcc cctaaccag gccctgcaca	3158
199	caaaggggca ggtgctggc tcagacctgc caagagccat atccggagg accctgcccc	3218
200	tgacctaaac ccacccaaa ggccaaactc tccactccct cagctggac accttctctc	3278

Input Set: I155514.RAW

Input Set: I155514.RAW

251	ctg cac aac cac tac acg cag aag agc ctc tcc ctg tct ccg ggt aaa	4225
252	Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys	
253	1165 1170 1175	
254	tga	4228

255	<210> 19	
256	<211> 4675	
257	<212> DNA	
258	<213> Homo sapien	
259	<220>	
260	<221> CDS	
261	<222> 1...3405, 3763...3807, 3926...4255, 4352...4675	
262	<300>	
263	<400> 19	
264	atg ggg cca gaa cgg aca ggg gcc gcg ccg ctg ccg ctg ctg gtg	48
265	Met Gly Pro Glu Arg Thr Gly Ala Ala Pro Leu Pro Leu Leu Val	
266	-25 -20 -15	
267	tta gcg ctc agt caa ggc att tta aat tgt tgt ttg gcc tac aat gtt	96
268	Leu Ala Leu Ser Gln Gly Ile Leu Asn Cys Cys Leu Ala Tyr Asn Val	
269	-10 -5 1	
270	ggt ctc cca gaa gca aaa ata ttt tcc ggt cct tca agt gaa cag ttt	144
271	Gly Leu Pro Glu Ala Lys Ile Phe Ser Gly Pro Ser Ser Glu Gln Phe	
272	5 10 15	
273	ggg tat gca gtg cag cag ttt ata aat cca aaa ggc aac tgg tta ctg	192
274	Gly Tyr Ala Val Gln Gln Phe Ile Asn Pro Lys Gly Asn Trp Leu Leu	
275	20 25 30 35	
276	gtt ggt tca ccc tgg agt ggc ttt cct gag aac cga atg gga gat gtg	240
277	Val Gly Ser Pro Trp Ser Gly Phe Pro Glu Asn Arg Met Gly Asp Val	
278	40 45 50	
279	tat aaa tgt cct gtt gac cta tcc act gcc aca tgt gaa aaa cta aat	288
280	Tyr Lys Cys Pro Val Asp Leu Ser Thr Ala Thr Cys Glu Lys Leu Asn	
281	55 60 65	
282	ttg caa act tca aca agc att cca aat gtt act gag atg aaa acc aac	336
283	Leu Gln Thr Ser Thr Ser Ile Pro Asn Val Thr Glu Met Lys Thr Asn	
284	70 75 80	
285	atg agc ctc ggc ttg atc ctc acc agg aac atg gga act gga ggt ttt	384
286	Met Ser Leu Gly Leu Ile Leu Thr Arg Asn Met Gly Thr Gly Phe	
287	85 90 95	
288	ctc aca tgt ggt cct ctg tgg gca cag caa tgt ggg aat cag tat tac	432
289	Leu Thr Cys Gly Pro Leu Trp Ala Gln Gln Cys Gly Asn Gln Tyr Tyr	
290	100 105 110 115	
291	aca acg ggt gtg tgt tct gac atc agt cct gat ttt cag ctc tca gcc	480
292	Thr Thr Gly Val Cys Ser Asp Ile Ser Pro Asp Phe Gln Leu Ser Ala	
293	120 125 130	
294	agc ttc tca cct gca act cag ccc tgc cct tcc ctc ata gat gtt gtg	528
295	Ser Phe Ser Pro Ala Thr Gln Pro Cys Pro Ser Leu Ile Asp Val Val	
296	135 140 145	
297	gtt gtg tgt gat gaa tca aat agt att tat cct tgg gat gca gta aag	576
298	Val Val Cys Asp Glu Ser Asn Ser Ile Tyr Pro Trp Asp Ala Val Lys	
299	150 155 160	
300	aat ttt ttg gaa aaa ttt gta caa ggc ctt gat ata ggc ccc aca aag	624

Input Set: I155514.RAW

301	Asn Phe Leu Glu Lys Phe Val Gln Gly Leu Asp Ile Gly Pro Thr Lys			
302	165	170	175	
303	aca cag gtg ggg tta att cag tat gcc aat aat cca aga gtt gtg ttt		672	
304	Thr Gln Val Gly Leu Ile Gln Tyr Ala Asn Asn Pro Arg Val Val Phe			
305	180	185	190	195
306	aac ttg aac aca tat aaa acc aaa gaa gaa atg att gta gca aca tcc		720	
307	Asn Leu Asn Thr Tyr Lys Thr Lys Glu Glu Met Ile Val Ala Thr Ser			
308	200	205	210	
309	cag aca tcc caa tat ggt ggg gac ctc aca aac aca ttc gga gca att		768	
310	Gln Thr Ser Gln Tyr Gly Asp Leu Thr Asn Thr Phe Gly Ala Ile			
311	215	220	225	
312	caa tat gca aga aaa tat gcc tat tca gca gct tct ggt ggg cga cga		816	
313	Gln Tyr Ala Arg Lys Tyr Ala Tyr Ser Ala Ala Ser Gly Gly Arg Arg			
314	230	235	240	
315	agt gct acg aaa gta atg gta gtt gta act gac ggt gaa tca cat gat		864	
316	Ser Ala Thr Lys Val Met Val Val Thr Asp Gly Glu Ser His Asp			
317	245	250	255	
318	ggt tca atg ttg aaa gct gtg att gat caa tgc aac cat gac aat ata		912	
319	Gly Ser Met Leu Lys Ala Val Ile Asp Gln Cys Asn His Asp Asn Ile			
320	260	265	270	275
321	ctg agg ttt ggc ata gca gtt ctt ggg tac tta aac aga aac gcc ctt		960	
322	Leu Arg Phe Gly Ile Ala Val Leu Gly Tyr Leu Asn Arg Asn Ala Leu			
323	280	285	290	
324	gat act aaa aat tta ata aaa gaa ata aaa gcg atc gct agt att cca		1008	
325	Asp Thr Lys Asn Leu Ile Lys Glu Ile Lys Ala Ile Ala Ser Ile Pro			
326	295	300	305	
327	aca gaa aga tac ttt ttc aat gtg tct gat gaa gca gct cta cta gaa		1056	
328	Thr Glu Arg Tyr Phe Phe Asn Val Ser Asp Glu Ala Ala Leu Leu Glu			
329	310	315	320	
330	aag gct ggg aca tta gga gaa caa att ttc agc att gaa ggt act gtt		1104	
331	Lys Ala Gly Thr Leu Gly Glu Gln Ile Phe Ser Ile Glu Gly Thr Val			
332	325	330	335	
333	caa gga gga gac aac ttt cag atg gaa atg tca caa gtg gga ttc agt		1152	
334	Gln Gly Gly Asp Asn Phe Gln Met Glu Met Ser Gln Val Gly Phe Ser			
335	340	345	350	355
336	gca gat tac tct tct caa aat gat att ctg atg ctg ggt gca gtg gga		1200	
337	Ala Asp Tyr Ser Ser Gln Asn Asp Ile Leu Met Leu Gly Ala Val Gly			
338	360	365	370	
339	gct ttt ggc tgg agt ggg acc att gtc cag aag aca tct cat ggc cat		1248	
340	Ala Phe Gly Trp Ser Gly Thr Ile Val Gln Lys Thr Ser His Gly His			
341	375	380	385	
342	ttg atc ttt cct aaa caa gcc ttt gac caa att ctg cag gac aga aat		1296	
343	Leu Ile Phe Pro Lys Gln Ala Phe Asp Gln Ile Leu Gln Asp Arg Asn			
344	390	395	400	
345	cac agt tca tat tta ggt tac tct gtg gct gca att tct act gga gaa		1344	
346	His Ser Ser Tyr Leu Gly Tyr Ser Val Ala Ala Ile Ser Thr Gly Glu			
347	405	410	415	
348	agc act cac ttt gtt gct ggt gct cct cgg gca aat tat acc ggc cag		1392	
349	Ser Thr His Phe Val Ala Gly Ala Pro Arg Ala Asn Tyr Thr Gly Gln			
350	420	425	430	435

Input Set: I155514.RAW

351	ata gtg cta tat agt gtg aat gag aat ggc aat atc acg gtt att cag	1440
352	Ile Val Leu Tyr Ser Val Asn Glu Asn Gly Asn Ile Thr Val Ile Gln	
353	440 445 450	
354	gct cac cga ggt gac cag att ggc tcc tat ttt ggt agt gtg ctg tgt	1488
355	Ala His Arg Gly Asp Gln Ile Gly Ser Tyr Phe Gly Ser Val Leu Cys	
356	455 460 465	
357	tca gtt gat gtg gat aaa gac acc att aca gac gtg ctc ttg gta ggt	1536
358	Ser Val Asp Val Asp Lys Asp Thr Ile Thr Asp Val Leu Leu Val Gly	
359	470 475 480	
360	gca cca atg tac atg agt gac cta aag aaa gag gaa gga aga gtc tac	1584
361	Ala Pro Met Tyr Met Ser Asp Leu Lys Lys Glu Glu Gly Arg Val Tyr	
362	485 490 495	
363	ctg ttt act atc aaa aag ggc att ttg ggt cag cac caa ttt ctt gaa	1632
364	Leu Phe Thr Ile Lys Lys Gly Ile Leu Gly Gln His Gln Phe Leu Glu	
365	500 505 510 515	
366	ggc ccc gag ggc att gaa aac act cga ttt ggt tca gca att gca gct	1680
367	Gly Pro Glu Gly Ile Glu Asn Thr Arg Phe Gly Ser Ala Ile Ala Ala	
368	520 525 530	
369	ctt tca gac atc aac atg gat ggc ttt aat gat gtg att gtt ggt tca	1728
370	Leu Ser Asp Ile Asn Met Asp Gly Phe Asn Asp Val Ile Val Gly Ser	
371	535 540 545	
372	cca cta gaa aat cag aat tct gga gct gta tac att tac aat ggt cat	1776
373	Pro Leu Glu Asn Gln Asn Ser Gly Ala Val Tyr Ile Tyr Asn Gly His	
374	550 555 560	
375	cag ggc act atc cgc aca aag tat tcc cag aaa atc ttg gga tcc gat	1824
376	Gln Gly Thr Ile Arg Thr Lys Tyr Ser Gln Lys Ile Leu Gly Ser Asp	
377	565 570 575	
378	gga gcc ttt agg agc cat ctc cag tac ttt ggg agg tcc ttg gat ggc	1872
379	Gly Ala Phe Arg Ser His Leu Gln Tyr Phe Gly Arg Ser Leu Asp Gly	
380	580 585 590 595	
381	tat gga gat tta aat ggg gat tcc atc acc gat gtg tct att ggt gcc	1920
382	Tyr Gly Asp Leu Asn Gly Asp Ser Ile Thr Asp Val Ser Ile Gly Ala	
383	600 605 610	
384	ttt gga caa gtg gtt caa ctc tgg tca caa agt att gct gat gta gct	1968
385	Phe Gly Gln Val Val Gln Leu Trp Ser Gln Ser Ile Ala Asp Val Ala	
386	615 620 625	
387	ata gaa gct tca ttc aca cca gaa aaa atc act ttg gtc aac aag aat	2016
388	Ile Glu Ala Ser Phe Thr Pro Glu Lys Ile Thr Leu Val Asn Lys Asn	
389	630 635 640	
390	gct cag ata att ctc aaa ctc tgc ttc agt gca aag ttc aga cct act	2064
391	Ala Gln Ile Ile Leu Lys Leu Cys Phe Ser Ala Lys Phe Arg Pro Thr	
392	645 650 655	
393	aag caa aac aat caa gtg gcc att gta tat aac atc aca ctt gat gca	2112
394	Lys Gln Asn Asn Gln Val Ala Ile Val Tyr Asn Ile Thr Leu Asp Ala	
395	660 665 670 675	
396	gat gga ttt tca tcc aga gta acc tcc agg ggg tta ttt aaa gaa aac	2160
397	Asp Gly Phe Ser Ser Arg Val Thr Ser Arg Gly Leu Phe Lys Glu Asn	
398	680 685 690	
399	aat gaa agg tgc ctg cag aag aat atg gta gta aat caa gca cag agt	2208
400	Asn Glu Arg Cys Leu Gln Lys Asn Met Val Val Asn Gln Ala Gln Ser	

Input Set: I155514.RAW

401	695	700	705	
402	tgc ccc gag cac atc att tat ata cag gag ccc tct gat gtt gtc aac			2256
403	Cys Pro Glu His Ile Ile Tyr Ile Gln Glu Pro Ser Asp Val Val Asn			
404	710	715	720	
405	tct ttg gat ttg cgt gtg gac atc agt ctg gaa aac cct ggc act agc			2304
406	Ser Leu Asp Leu Arg Val Asp Ile Ser Leu Glu Asn Pro Gly Thr Ser			
407	725	730	735	
408	cct gcc ctt gaa gcc tat tct gag act gcc aag gtc ttc agt att cct			2352
409	Pro Ala Leu Glu Ala Tyr Ser Glu Thr Ala Lys Val Phe Ser Ile Pro			
410	740	745	750	755
411	ttc cac aaa gac tgt ggt gag gat gga ctt tgc att tct gat cta gtc			2400
412	Phe His Lys Asp Cys Gly Glu Asp Gly Leu Cys Ile Ser Asp Leu Val			
413	760	765	770	
414	cta gat gtc cga caa ata cca gct gct caa gaa caa ccc ttt att gtc			2448
415	Leu Asp Val Arg Gln Ile Pro Ala Ala Gln Glu Gln Pro Phe Ile Val			
416	775	780	785	
417	agc aac caa aac aaa agg tta aca ttt tca gta aca ctg aaa aat aaa			2496
418	Ser Asn Gln Asn Lys Arg Leu Thr Phe Ser Val Thr Leu Lys Asn Lys			
419	790	795	800	
420	agg gaa agt gca tac aac act gga att gtt gtt gat ttt tca gaa aac			2544
421	Arg Glu Ser Ala Tyr Asn Thr Gly Ile Val Val Asp Phe Ser Glu Asn			
422	805	810	815	
423	ttg ttt ttt gca tca ttc tcc cta ccg gtt gat ggg aca gaa gta aca			2592
424	Leu Phe Ala Ser Phe Ser Leu Pro Val Asp Gly Thr Glu Val Thr			
425	820	825	830	835
426	tgc cag gtg gct gca tct cag aag tct gtt gcc tgc gat gta ggc tac			2640
427	Cys Gln Val Ala Ala Ser Gln Lys Ser Val Ala Cys Asp Val Gly Tyr			
428	840	845	850	
429	cct gct tta aag aga gaa caa cag gtg act ttt act att aac ttt gac			2688
430	Pro Ala Leu Lys Arg Glu Gln Gln Val Thr Phe Thr Ile Asn Phe Asp			
431	855	860	865	
432	ttc aat ctt caa aac ctt cag aat cag gcg tct ctc agt ttc caa gcc			2736
433	Phe Asn Leu Gln Asn Leu Gln Asn Gln Ala Ser Leu Ser Phe Gln Ala			
434	870	875	880	
435	tta agt gaa agc caa gaa aac aag gct gat aat ttg gtc aac ctc			2784
436	Leu Ser Glu Ser Gln Glu Glu Asn Lys Ala Asp Asn Leu Val Asn Leu			
437	885	890	895	
438	aaa att cct ctc ctg tat gat gct gaa att cac tta aca aga tct acc			2832
439	Lys Ile Pro Leu Leu Tyr Asp Ala Glu Ile His Leu Thr Arg Ser Thr			
440	900	905	910	915
441	aac ata aat ttt tat gaa atc tct tcg gat ggg aat gtt cct tca atc			2880
442	Asn Ile Asn Phe Tyr Glu Ile Ser Ser Asp Gly Asn Val Pro Ser Ile			
443	920	925	930	
444	gtg cac agt ttt gaa gat gtt ggt cca aaa ttc atc ttc tcc ctg aag			2928
445	Val His Ser Phe Glu Asp Val Gly Pro Lys Phe Ile Phe Ser Leu Lys			
446	935	940	945	
447	gta aca aca gga agt gtt cca gta agc atg gca act gta atc atc cac			2976
448	Val Thr Thr Gly Ser Val Pro Val Ser Met Ala Thr Val Ile Ile His			
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Input Set: I155514.RAW

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 452 965 970 975
 453 gtg caa aca gac aag gct ggt gac atc agt tgt aat gca gat atc aat 3072
 454 Val Gln Thr Asp Lys Ala Gly Asp Ile Ser Cys Asn Ala Asp Ile Asn
 455 980 985 990 995
 456 cca ctg aaa ata gga caa aca tct tct tct gta tct ttc aaa agt gaa 3120
 457 Pro Leu Lys Ile Gly Gln Thr Ser Ser Val Ser Phe Lys Ser Glu
 458 1000 1005 1010
 459 aat ttc agg cac acc aaa gaa ttg aac tgc aga act gct tcc tgt agt 3168
 460 Asn Phe Arg His Thr Lys Glu Leu Asn Cys Arg Thr Ala Ser Cys Ser
 461 1015 1020 1025
 462 aat gtt acc tgc tgg ttg aaa gac gtt cac atg aaa gga gaa tac ttt 3216
 463 Asn Val Thr Cys Trp Leu Lys Asp Val His Met Lys Gly Glu Tyr Phe
 464 1030 1035 1040
 465 gtt aat gtg act acc aga att tgg aac ggg act ttc gca tca tca acg 3264
 466 Val Asn Val Thr Thr Arg Ile Trp Asn Gly Thr Phe Ala Ser Ser Thr
 467 1045 1050 1055
 468 ttc cag aca gta cag cta acg gca gct gca gaa atc aac acc tat aac 3312
 469 Phe Gln Thr Val Gln Leu Thr Ala Ala Ala Glu Ile Asn Thr Tyr Asn
 470 1060 1065 1070 1075
 471 cct gag ata tat gtg att gaa gat aac act gtt acg att ccc ctg atg 3360
 472 Pro Glu Ile Tyr Val Ile Glu Asp Asn Thr Val Thr Ile Pro Leu Met
 473 1080 1085 1090
 474 ata atg aaa cct gat gag aaa gcc gaa gta cca aca gat ccc gag 3405
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 483 gag ccc aaa tct tgt gac aaa act cac aca tgc cca ccg tgc cca 3807
 484 Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro
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 487 tgcattccagg gacaggcccc agccgggtgc tgacacgtcc acctccatct cttccctca 3925
 488 gca cct gaa ctc ctg ggg gga ccg tca gtc ttc ctc ttc ccc cca aaa 3973
 489 Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys
 490 1125 1130 1135
 491 ccc aag gac acc ctc atg atc tcc cgg acc cct gag gtc aca tgc gtg 4021
 492 Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val
 493 1140 1145 1150
 494 gtg gtg gac gtg agc cac gaa gac cct gag gtc aag ttc aac tgg tac 4069
 495 Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr
 496 1155 1160 1165
 497 gtg gac ggc gtg gag gtg cat aat gcc aag aca aag ccg cgg gag gag 4117
 498 Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu
 499 1170 1175 1180 1185
 500 cag tac aac agc acg tac cgg gtg gtc agc gtc ctc acc gtc ctg cac 4165

Input Set: I155514.RAW

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502 1190 1195 1200
503 cag gac tgg ctg aat ggc aag gag tac aag tgc aag gtc tcc aac aaa 4213
504 Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys
505 1205 1210 1215
506 gcc ctc cca gcc ccc atc gag aaa acc atc tcc aaa gcc aaa 4255
507 Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys
508 1220 1225 1230
509 ggtgggaccc gtgggggtgcg agggccacat ggacagaggc cggctcgcc caccctctgc 4315
510 cctgagatg accgctgtac caacctctgt cctaca ggg cag ccc cga gaa cca 4369
511 Gly Gln Pro Arg Glu Pro
512 1235
513 cag gtg tac acc ctg ccc cca tcc cgg gat gag ctg acc aag aac cag 4417
514 Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln
515 1240 1245 1250
516 gtc agc ctg acc tgc ctg gtc aaa ggc ttc tat ccc agc gac atc gcc 4465
517 Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala
518 1255 1260 1265
519 gtg gag tgg gag agc aat ggg cag ccg gag aac aac tac aag acc acg 4513
520 Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr
521 1270 1275 1280 1285
522 cct ccc gtg ctg gat tcc gac ggc tcc ttc ttc ctc tac agc aag ctc 4561
523 Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu
524 1290 1295 1300
525 acc gtg gac aag agc agg tgg cag cag ggg aac gtc ttc tca tgc tcc 4609
526 Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser
527 1305 1310 1315
528 gtg atg cat gag gct ctg cac aac cac tac acg cag aag agc ctc tcc 4657
529 Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser
530 1320 1325 1330
531 ctg tct ccg ggt aaa tga 4675
532 Leu Ser Pro Gly Lys
533 1335

PAGE: 1

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/155,514DATE: 02/03/2000
TIME: 13:10:50

Input Set: I155514.RAW

This Raw Listing contains the General
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containing ERRORS.

Does Not Comply
Corrected Diskette Needed

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2 Tanaka, Toshiaki
3 <120> Chimeric proteins, their heterodimer complexes, and platelet
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27 ggc ccc gaa gct gca ctc cgg gag acg gtg atg ctg ttg ctg tgc ctg 96
28 Gly Pro Glu Ala Ala Leu Arg Glu Thr Val Met Leu Leu Leu Cys Leu
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31 Gly Val Pro Thr Gly Arg Pro Tyr Asn Val Asp Thr Glu Ser Ala Leu
32 -5 1 5
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34 Leu Tyr Gln Gly Pro His Asn Thr Leu Phe Gly Tyr Ser Val Val Leu
35 10 15 20 25
36 cac agc cac ggg gcg aac cga tgg ctc cta gtg ggt gcg ccc act gcc 240
37 His Ser His Gly Ala Asn Arg Trp Leu Leu Val Gly Ala Pro Thr Ala
38 30 35 40
39 aac tgg ctc gcc aac gct tca gtg atc aat ccc ggg gcg att tac aga 288

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PAGE: 2

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/155,514DATE: 02/03/2000
TIME: 13:10:50

Input Set: I155514.RAW

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43	Cys Arg Ile Gly Lys Asn Pro Gly Gln Thr Cys Glu Gln Leu Gln Leu			
44	60	65	70	
45	ggt agc cct aat gga gaa cct tgt gga aag act tgt ttg gaa gag aga	384		
46	Gly Ser Pro Asn Gly Glu Pro Cys Gly Lys Thr Cys Leu Glu Glu Arg			
47	75	80	85	
48	gac aat cag tgg ttg ggg gtc aca ctt tcc aga cag cca gga gaa aat	432		
49	Asp Asn Gln Trp Leu Gly Val Thr Leu Ser Arg Gln Pro Gly Glu Asn			
50	90	95	100	105
51	gga tcc atc gtg act tgt ggg cat aga tgg aaa aat ata ttt tac ata	480		
52	Gly Ser Ile Val Thr Cys Gly His Arg Trp Lys Asn Ile Phe Tyr Ile			
53	110	115	120	
54	aag aat gaa aat aag ctc ccc act ggt ggt tgc tat gga gtg ccc cct	528		
55	Lys Asn Glu Asn Lys Leu Pro Thr Gly Gly Cys Tyr Gly Val Pro Pro			
56	125	130	135	
57	gat tta cga aca gaa ctg agt aaa aga ata gct ccg tgt tat caa gat	576		
58	Asp Leu Arg Thr Glu Leu Ser Lys Arg Ile Ala Pro Cys Tyr Gln Asp			
59	140	145	150	
60	tat gtg aaa aaa ttt gga gaa aat ttt gca tca tgt caa gct gga ata	624		
61	Tyr Val Lys Lys Phe Gly Glu Asn Phe Ala Ser Cys Gln Ala Gly Ile			
62	155	160	165	
63	tcc agt ttt tac aca aag gat tta att gtg atg ggg gcc cca gga tca	672		
64	Ser Ser Phe Tyr Thr Lys Asp Leu Ile Val Met Gly Ala Pro Gly Ser			
65	170	175	180	185
66	tct tac tgg act ggc tct ctt ttt gtc tac aat ata act aca aat aaa	720		
67	Ser Tyr Trp Thr Gly Ser Leu Phe Val Tyr Asn Ile Thr Thr Asn Lys			
68	190	195	200	
69	tac aag gct ttt tta gac aaa caa aat caa gta aaa ttt gga agt tat	768		
70	Tyr Lys Ala Phe Leu Asp Lys Gln Asn Gln Val Lys Phe Gly Ser Tyr			
71	205	210	215	
72	tta gga tat tca gtc gga gct ggt cat ttt cgg agc cag cat act acc	816		
73	Leu Gly Tyr Ser Val Gly Ala Gly His Phe Arg Ser Gln His Thr Thr			
74	220	225	230	
75	gaa gta gtc gga gga gct cct caa cat gag cag att ggt aag gca tat	864		
76	Glu Val Val Gly Gly Ala Pro Gln His Glu Gln Ile Gly Lys Ala Tyr			
77	235	240	245	
78	ata ttc agc att gat gaa aaa gaa cta aat atc tta cat gaa atg aaa	912		
79	Ile Phe Ser Ile Asp Glu Lys Glu Leu Asn Ile Leu His Glu Met Lys			
80	250	255	260	265
81	ggt aaa aag ctt gga tcg tac ttt gga gct tct gtc tgt gct gtg gac	960		
82	Gly Lys Lys Leu Gly Ser Tyr Phe Gly Ala Ser Val Cys Ala Val Asp			
83	270	275	280	
84	ctc aat gca gat ggc ttc tca gat ctg ctc gtg gga gca ccc atg cag	1008		
85	Leu Asn Ala Asp Gly Phe Ser Asp Leu Leu Val Gly Ala Pro Met Gln			
86	285	290	295	
87	agc acc atc aga gag gaa gga aga gtg ttt gtg tac atc aac tct ggc	1056		
88	Ser Thr Ile Arg Glu Glu Gly Arg Val Phe Val Tyr Ile Asn Ser Gly			
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PAGE: 3

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/155,514DATE: 02/03/2000
TIME: 13:10:50

Input Set: I155514.RAW

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92	315 320 325	
93	aaa tat gct gca aga ttt ggg gaa tct ata gtt aat ctt ggc gac att	1152
94	Lys Tyr Ala Ala Arg Phe Gly Glu Ser Ile Val Asn Leu Gly Asp Ile	
95	330 335 340 345	
96	gac aat gat ggc ttt gaa gat gtt gct atc gga gct cca caa gaa gat	1200
97	Asp Asn Asp Gly Phe Glu Asp Val Ala Ile Gly Ala Pro Gln Glu Asp	
98	350 355 360	
99	gac ttg caa ggt gct att tat att tac aat ggc cgt gca gat ggg atc	1248
100	Asp Leu Gln Gly Ala Ile Tyr Ile Tyr Asn Gly Arg Ala Asp Gly Ile	
101	365 370 375	
102	tcg tca acc ttc tca cag aga att gaa gga ctt cag atc agc aaa tcg	1296
103	Ser Ser Thr Phe Ser Gln Arg Ile Glu Gly Leu Gln Ile Ser Lys Ser	
104	380 385 390	
105	tta agt atg ttt gga cag tct ata tca gga caa att gat gca gat att	1344
106	Leu Ser Met Phe Gly Gln Ser Ile Ser Gly Gln Ile Asp Ala Asp Asn	
107	395 400 405	
108	aat ggc tat gta gat gta gca ggt cgt gct ttt cgg tct gat tct gct	1392
109	Asn Gly Tyr Val Asp Val Ala Val Gly Ala Phe Arg Ser Asp Ser Ala	
110	410 415 420 425	
111	gtc ttg cta agg aca aga cct gta gta att gtt gac gct tct tta agc	1440
112	Val Leu Leu Arg Thr Arg Pro Val Val Ile Val Asp Ala Ser Leu Ser	
113	430 435 440	
114	cac cct gag tca gta aat aga acg aaa ttt gac tgt gtt gaa aat gga	1488
115	His Pro Glu Ser Val Asn Arg Thr Lys Phe Asp Cys Val Glu Asn Gly	
116	445 450 455	
117	tgg cct tct gtg tgc ata gat cta aca ctt tgt ttc tca tat aag ggc	1536
118	Trp Pro Ser Val Cys Ile Asp Leu Thr Leu Cys Phe Ser Tyr Lys Gly	
119	460 465 470	
120	aag gaa gtt cca ggt tac att gtt ttg ttt tat aac atg agt ttg gat	1584
121	Lys Glu Val Pro Gly Tyr Ile Val Leu Phe Tyr Asn Met Ser Leu Asp	
122	475 480 485	
123	gtg aac aga aag gca gag tct cca cca aga ttc tat ttc tct tct aat	1632
124	Val Asn Arg Lys Ala Glu Ser Pro Pro Arg Phe Tyr Phe Ser Ser Asn	
125	490 495 500 505	
126	gga act tct gac gtg att aca gga agc ata cag gtg tcc agc aga gaa	1680
127	Gly Thr Ser Asp Val Ile Thr Gly Ser Ile Gln Val Ser Ser Arg Glu	
128	510 515 520	
129	gct aac tgt aga aca cat caa gca ttt atg cgg aaa gat gtg cgg gac	1728
130	Ala Asn Cys Arg Thr His Gln Ala Phe Met Arg Lys Asp Val Arg Asp	
131	525 530 535	
132	atc ctc acc cca att cag att gaa gct gct tac cac ctt ggt cct cat	1776
133	Ile Leu Thr Pro Ile Gln Ile Glu Ala Ala Tyr His Leu Gly Pro His	
134	540 545 550	
135	gtc atc agt aaa cga agt aca gag gaa ttc cca cca ott cag cca att	1824
136	Val Ile Ser Lys Arg Ser Thr Glu Glu Phe Pro Pro Leu Gln Pro Ile	
137	555 560 565	
138	ctt cag cag aag aaa gaa aaa gac ata atg aaa aaa aca ata aac ttt	1872
139	Leu Gln Gln Lys Lys Glu Lys Asp Ile Met Lys Lys Thr Ile Asn Phe	

PAGE: 4

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/155,514DATE: 02/03/2000
TIME: 13:10:50

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143	590	595	600		
144	gca aag att ggg ttt ttg aag ccc cat gaa aat aaa aca tat ctt gct				1968
145	Ala Lys Ile Gly Phe Leu Lys Pro His Glu Asn Lys Thr Tyr Leu Ala				
146	605	610	615		
147	gtt ggg agt atg aag aca ttg atg ttg aat gtg tcc ttg ttt aat gct				2016
148	Val Gly Ser Met Lys Thr Leu Met Leu Asn Val Ser Leu Phe Asn Ala				
149	620	625	630		
150	gga gat gat gca tat gaa acg act cta cat gtc aaa cta ccc gtg ggt				2064
151	Gly Asp Asp Ala Tyr Glu Thr Thr Leu His Val Lys Leu Pro Val Gly				
152	635	640	645		
153	ctt tat ttc att aag att tta gag ctg gaa gag aag caa ata aac tgt				2112
154	Leu Tyr Phe Ile Lys Ile Leu Glu Leu Glu Lys Gln Ile Asn Cys				
155	650	655	660	665	
156	gaa gtc aca gat aac tct ggc gtg gta caa ctt gac tgc agt att ggc				2160
157	Glu Val Thr Asp Asn Ser Gly Val Val Gln Leu Asp Cys Ser Ile Gly				
158	670	675	680		
159	tat ata tat gta gat cat ctc tca agg ata gat att agc ttt ctc ctg				2208
160	Tyr Ile Tyr Val Asp His Leu Ser Arg Ile Asp Ile Ser Phe Leu Leu				
161	685	690	695		
162	gat gtg agc tca ctc agc aga gcg gaa gag gac ctc agt atc aca gtg				2256
163	Asp Val Ser Ser Leu Ser Arg Ala Glu Glu Asp Leu Ser Ile Thr Val				
164	700	705	710		
165	cat gct acc tgt gaa aat gaa gag gaa atg gac aat cta aag cac agc				2304
166	His Ala Thr Cys Glu Asn Glu Glu Met Asp Asn Leu Lys His Ser				
167	715	720	725		
168	aga gtg act gta gca ata cct tta aaa tat gag gtt aag ctg act gtt				2352
169	Arg Val Thr Val Ala Ile Pro Leu Lys Tyr Glu Val Lys Leu Thr Val				
170	730	735	740	745	
171	cat ggg ttt gta aac cca act tca ttt gtg tat gga tca aat gat gaa				2400
172	His Gly Phe Val Asn Pro Thr Ser Phe Val Tyr Gly Ser Asn Asp Glu				
173	750	755	760		
174	aat gag cct gaa acg tgc atg gtg gag aaa atg aac tta act ttc cat				2448
175	Asn Glu Pro Glu Thr Cys Met Val Glu Lys Met Asn Leu Thr Phe His				
176	765	770	775		
177	gtt atc aac act ggc aat agt atg gct ccc aat gtt agt gtg gaa ata				2496
178	Val Ile Asn Thr Gly Asn Ser Met Ala Pro Asn Val Ser Val Glu Ile				
179	780	785	790		
E--> 180	atg gta cca aat tct ttt agc ccc caa act gat aag ctg ttc aac att				2588
181	Met Val Pro Asn Ser Phe Ser Pro Gln Thr Asp Lys Leu Phe Asn Ile				2544
182	795	800	805		
183	ttg gat gtc cag act act gga gaa tgc cac ttt gaa aat tat caa				2592
184	Leu Asp Val Gln Thr Thr Gly Glu Cys His Phe Glu Asn Tyr Gln				
185	810	815	820	825	
186	aga gtg tgt gca tta gag cag caa aag agt gca atg cag acc ttg aaa				2640
187	Arg Val Cys Ala Leu Glu Gln Gln Lys Ser Ala Met Gln Thr Leu Lys				
188	830	835	840		
189	ggc ata gtc cgg ttc ttg tcc aag act gat aag agg cta ttg tac tgc				2688

PAGE: 5

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/155,514DATE: 02/03/2000
TIME: 13:10:50

Input Set: I155514.RAW

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 192 ata aaa gct gat cca cat tgt tta aat ttc ttg tgt aat ttt ggg aaa 2736
 193 Ile Lys Ala Asp Pro His Cys Leu Asn Phe Leu Cys Asn Phe Gly Lys
 194 860 865 870
 195 atg gaa agt gga aaa gaa gcc agt gtt cat atc caa ctg gaa ggc cg 2784
 196 Met Glu Ser Gly Lys Glu Ala Ser Val His Ile Gln Leu Glu Gly Arg
 197 875 880 885
 198 cca tcc att tta gaa atg gat gag act tca gca ctc aag ttt gaa ata 2832
 199 Pro Ser Ile Leu Glu Met Asp Glu Thr Ser Ala Leu Lys Phe Glu Ile
 200 890 895 900 905
 201 aga gca aca ggt ttt cca gag cca aat cca aga gta att gaa cta aac 2880
 202 Arg Ala Thr Gly Phe Pro Glu Pro Asn Pro Arg Val Ile Glu Leu Asn
 203 910 915 920
 204 aag gat gag aat gtt gcg cat gtt cta ctg gaa gga cta cat cat caa 2928
 205 Lys Asp Glu Asn Val Ala His Val Leu Leu Glu Gly Leu His His Gln
 206 925 930 935
 207 aga ccc aaa cgt tat ttc acg gat ccc gag ctgctggaag caggctcagc 2978
 208 Arg Pro Lys Arg Tyr Phe Thr Asp Pro Glu
 209 940 945
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 219 Lys Thr His Thr Cys Pro Pro Cys Pro
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 223 Ala Pro Glu Leu Leu
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 226 Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu
 227 970 975 980
 228 atg atc tcc cgg acc cct gag gtc aca tgc gtg gtg gac gtg agc 3589
 229 Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser
 230 985 990 995
 231 cac gaa gac cct gag gtc aag ttc aac tgg tac gtg gac ggc gtg gag 3637
 232 His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu
 233 1000 1005 1010 1015
 234 gtg cat aat gcc aag aca aag ccg cgg gag gag cag tac aac agc acg 3685
 235 Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr
 236 1020 1025 1030
 237 tac cgg gtg gtc agc gtc ctc acc gtc ctg cac cag gac tgg ctg aat 3733
 238 Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn
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PAGE: 6

RAW SEQUENCE LISTING
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TIME: 13:10:50

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 243 atc gag aaa acc atc tcc aaa gcc aaa ggtgggaccc gtgggggtgcg 3828
 244 Ile Glu Lys Thr Ile Ser Lys Ala Lys
 245 1065 1070
 246 agggccacat ggacagaggg cggctcgcc caccctctgc cctgagagtg accgctgtac 3888
 247 caaccctctgt cctaca ggg cag ccc cga gaa cca cag gtg tac acc ctg 3937
 248 Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu
 249 1075 1080
 250 ccc cca tcc cgg gat gag ctg acc aag aac cag gtc agc ctg acc tgc 3985
 251 Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val Ser Leu Thr Cys
 252 1085 1090 1095
 253 ctg gtc aaa ggc ttc tat ccc agc gac atc gcc gtg gag tgg gag agc 4033
 254 Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser
 255 1100 1105 1110 1115
 256 aat ggg cag ccg gag aac aac tac aag acc acg cct ccc gtg ctg gat 4081
 257 Asn Gly Gln Pro Glu Asn Asn Tyr Thr Thr Pro Pro Val Leu Asp
 258 1120 1125 1130
 259 tcc gac ggc tcc ttc ctc tac agc aag ctc acc gtg gac aag agc 4129
 260 Ser Asp Gly Ser Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser
 261 1135 1140 1145
 262 agg tgg cag cag ggg aac gtc ttc tca tgc tcc gtg atg cat gag gct 4177
 263 Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala
 264 1150 1155 1160
 265 ctg cac aac cac tac acg cag aag agc ctc tcc ctg tct ccg ggt aaa 4225
 266 Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Pro Gly Lys
 267 1165 1170 1175
 268 tga 4228

269 <210> 19
 270 <211> 4675
 271 <212> DNA
 272 <213> Homo sapien
 273 <220>
 274 <221> CDS
 275 <222> 1...3405, 3763...3807, 3926...4255, 4352...4675
 276 <300>
 277 <400> 19
 278 atg ggg cca gaa cgg aca ggg gcc gcg ccg ctg ccg ctg ctg gtg 48
 279 Met Gly Pro Glu Arg Thr Gly Ala Ala Pro Leu Pro Leu Leu Val
 280 -25 -20 -15
 281 tta gcg ctc agt caa ggc att tta aat tgt tgt ttg gcc tac aat gtt 96
 282 Leu Ala Leu Ser Gln Gly Ile Leu Asn Cys Cys Leu Ala Tyr Asn Val
 283 -10 -5 1
 284 ggt ctc cca gaa gca aaa ata ttt tcc ggt cct tca agt gaa cag ttt
 285 Gly Leu Pro Glu Ala Lys Ile Phe Ser Gly Pro Ser Ser Glu Gln Phe
 286 5 10 15
 287 ggg tat gca gtg cag cag ttt ata aat cca aaa ggc aac tgg tta ctg
 288 Gly Tyr Ala Val Gln Gln Phe Ile Asn Pro Lys Gly Asn Trp Leu Leu 192

(114) 144

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PAGE: 7

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/155,514DATE: 02/03/2000
TIME: 13:10:50

Input Set: I155514.RAW

289	20	25	30	35	
290	gtt ggt tca ccc tgg agt ggc ttt cct gag aac cga atg gga gat gtg				240
291	Val Gly Ser Pro Trp Ser Gly Phe Pro Glu Asn Arg Met Gly Asp Val				
292	40	45	50		
293	tat aaa tgt cct gtt gac cta tcc act gcc aca tgt gaa aaa cta aat				288
294	Tyr Lys Cys Pro Val Asp Leu Ser Thr Ala Thr Cys Glu Lys Leu Asn				
295	55	60	65		
296	ttg caa act tca aca agc att cca aat gtt act gag atg aaa acc aac				336
297	Leu Gln Thr Ser Thr Ser Ile Pro Asn Val Thr Glu Met Lys Thr Asn				
298	70	75	80		
299	atg agc ctc ggc ttg atc ctc acc agg aac atg gga act gga ggt ttt				384
300	Met Ser Leu Gly Leu Ile Leu Thr Arg Asn Met Gly Thr Gly Phe				
301	85	90	95		
302	ctc aca tgt ggt cct ctg tgg gca cag caa tgt ggg aat cag tat tac				432
303	Leu Thr Cys Gly Pro Leu Trp Ala Gln Gln Cys Gly Asn Gln Tyr Tyr				
304	100	105	110	115	
305	aca acg ggt gtg tgt tct gac atc agt cct gat ttt cag ctc tca gcc				480
306	Thr Thr Gly Val Cys Ser Asp Ile Ser Pro Asp Phe Gln Leu Ser Ala				
307	120	125	130		
308	agc ttc tca cct gca act cag ccc tgc cct tcc ctc ata gat gtt gtg				528
309	Ser Phe Ser Pro Ala Thr Gln Pro Cys Pro Ser Leu Ile Asp Val Val				
310	135	140	145		
311	gtt gtg tgt gat gaa tca aat agt att tat cct tgg gat gca gta aag				576
312	Val Val Cys Asp Glu Ser Asn Ser Ile Tyr Pro Trp Asp Ala Val Lys				
313	150	155	160		
314	aat ttt ttg gaa aaa ttt gta caa ggc ctt gat ata ggc ccc aca aag				624
315	Asn Phe Leu Glu Lys Phe Val Gln Gly Leu Asp Ile Gly Pro Thr Lys				
316	165	170	175		
317	aca cag gtg ggg tta att cag tat gcc aat aat cca aga gtt gtg ttt				672
318	Thr Gln Val Gly Leu Ile Gln Tyr Ala Asn Asn Pro Arg Val Val Phe				
319	180	185	190	195	
320	aac ttg aac aca tat aaa acc aaa gaa gaa atg att gta gca aca tcc				720
321	Asn Leu Asn Thr Tyr Lys Thr Lys Glu Glu Met Ile Val Ala Thr Ser				
322	200	205	210		
323	cag aca tcc caa tat ggt ggg gac ctc aca aac aca ttc gga gca att				768
324	Gln Thr Ser Gln Tyr Gly Gly Asp Leu Thr Asn Thr Phe Gly Ala Ile				
325	215	220	225		
326	caa tat gca aga aaa tat gcc tat tca gca gct tct ggt ggg cga cga				816
327	Gln Tyr Ala Arg Lys Tyr Ala Tyr Ser Ala Ala Ser Gly Gly Arg Arg				
328	230	235	240		
329	agt gct acg aaa gta atg gta gtt gta act gac ggt gaa tca cat gat				864
330	Ser Ala Thr Lys Val Met Val Val Val Thr Asp Gly Glu Ser His Asp				
331	245	250	255		
332	ggt tca atg ttg aaa gct gtg att gat caa tgc aac cat gac aat ata				912
333	Gly Ser Met Leu Lys Ala Val Ile Asp Gln Cys Asn His Asp Asn Ile				
334	260	265	270	275	
335	ctg agg ttt ggc ata gca gtt ctt ggg tac tta aac aga aac gcc ctt				960
336	Leu Arg Phe Gly Ile Ala Val Leu Gly Tyr Leu Asn Arg Asn Ala Leu				
337	280	285	290		
338	gat act aaa aat tta ata aaa gaa ata aaa gcg atc gct agt att cca				1008

PAGE: 8

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/155,514DATE: 02/03/2000
TIME: 13:10:50

Input Set: I155514.RAW

339	Asp Thr Lys Asn Leu Ile Lys Glu Ile Lys Ala Ile Ala Ser Ile Pro			
340	295	300	305	
341	aca gaa aga tac ttt ttc aat gtg tct gat gaa gca gct cta cta gaa		1056	
342	Thr Glu Arg Tyr Phe Phe Asn Val Ser Asp Glu Ala Ala Leu Leu Glu			
343	310	315	320	
344	aag gct ggg aca tta gga gaa caa att ttc agc att gaa ggt act gtt		1104	
345	Lys Ala Gly Thr Leu Gly Glu Gln Ile Phe Ser Ile Glu Gly Thr Val			
346	325	330	335	
347	caa gga gga gac aac ttt cag atg gaa atg tca caa gtg gga ttc agt		1152	
348	Gln Gly Gly Asp Asn Phe Gln Met Glu Met Ser Gln Val Gly Phe Ser			
349	340	345	350	355
350	gca gat tac tct tct caa aat gat att ctg atg ctg ggt gca gtg gga		1200	
351	Ala Asp Tyr Ser Ser Gln Asn Asp Ile Leu Met Leu Gly Ala Val Gly			
352	360	365	370	
353	gct ttt ggc tgg agt ggg acc att gtc cag aag aca tct cat ggc cat		1248	
354	Ala Phe Gly Trp Ser Gly Thr Ile Val Gln Lys Thr Ser His Gly His			
355	375	380	385	
356	ttg atc ttt cct aaa caa gcc ttt gac caa att ctg cag gac aga aat		1296	
357	Leu Ile Phe Pro Lys Gln Ala Phe Asp Gln Ile Leu Gln Asp Arg Asn			
358	390	395	400	
359	cac agt tca tat tta ggt tac tct gtg gct gca att tct act gga gaa		1344	
360	His Ser Ser Tyr Leu Gly Tyr Ser Val Ala Ala Ile Ser Thr Gly Glu			
361	405	410	415	
362	agc act cac ttt gtt ggt gct gct cct cgg gca aat tat acc ggc cag		1392	
363	Ser Thr His Phe Val Ala Gly Ala Pro Arg Ala Asn Tyr Thr Gly Gln			
364	420	425	430	435
365	ata gtg cta tat agt gtg aat gag aat ggc aat atc acg gtt att cag		1440	
366	Ile Val Leu Tyr Ser Val Asn Glu Asn Ile Thr Val Ile Gln			
367	440	445	450	
368	gct cac cga ggt gac cag att ggc tcc tat ttt ggt agt gtg ctg tgt		1488	
369	Ala His Arg Gly Asp Gln Ile Gly Ser Tyr Phe Gly Ser Val Leu Cys			
370	455	460	465	
371	tca gtt gat gtg gat aaa gac acc att aca gac gtg ctc ttg gta ggt		1536	
372	Ser Val Asp Val Asp Lys Asp Thr Ile Thr Asp Val Leu Leu Val Gly			
373	470	475	480	
374	gca cca atg tac atg agt gac cta aag aaa gag gaa gga aga gtc tac		1584	
375	Ala Pro Met Tyr Met Ser Asp Leu Lys Glu Gly Arg Val Tyr			
376	485	490	495	
377	ctg ttt act atc aaa aag ggc att ttg ggt cag cac caa ttt ctt gaa		1632	
378	Leu Phe Thr Ile Lys Lys Gly Ile Leu Gly Gln His Gln Phe Leu Glu			
379	500	505	510	515
380	ggc ccc gag ggc att gaa aac act cga ttt ggt tca gca att gca gct		1680	
381	Gly Pro Glu Gly Ile Glu Asn Thr Arg Phe Gly Ser Ala Ile Ala Ala			
382	520	525	530	
383	ctt tca gac atc aac atg gat ggc ttt aat gat gtg att gtt ggt tca		1728	
384	Leu Ser Asp Ile Asn Met Asp Gly Phe Asn Asp Val Ile Val Gly Ser			
385	535	540	545	
386	cca cta gaa aat cag aat tct gga gct gta tac att tac aat ggt cat		1776	
387	Pro Leu Glu Asn Gln Asn Ser Gly Ala Val Tyr Ile Tyr Asn Gly His			
388	550	555	560	

PAGE: 9

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/155,514DATE: 02/03/2000
TIME: 13:10:50

Input Set: I155514.RAW

389	cag ggc act atc cgc aca aag tat tcc cag aaa atc ttg gga tcc gat	1824
390	Gln Gly Thr Ile Arg Thr Lys Tyr Ser Gln Lys Ile Leu Gly Ser Asp	
391	565 570 575	
392	gga gcc ttt agg agc cat ctc cag tac ttt ggg agg tcc ttg gat ggc	1872
393	Gly Ala Phe Arg Ser His Leu Gln Tyr Phe Gly Arg Ser Leu Asp Gly	
394	580 585 590 595	
395	tat gga gat tta aat ggg gat tcc atc acc gat gtg tct att ggt gcc	1920
396	Tyr Gly Asp Leu Asn Gly Asp Ser Ile Thr Asp Val Ser Ile Gly Ala	
397	600 605 610	
398	ttt gga caa gtg gtt caa ctc tgg tca caa agt att gct gat gta gct	1968
399	Phe Gly Gln Val Val Gln Leu Trp Ser Gln Ser Ile Ala Asp Val Ala	
400	615 620 625	
401	ata gaa gct tca ttc aca cca gaa aaa atc act ttg gtc aac aag aat	2016
402	Ile Glu Ala Ser Phe Thr Pro Glu Lys Ile Thr Leu Val Asn Lys Asn	
403	630 635 640	
404	gct cag ata att ctc aaa ctc tgc ttc agt gca aag ttc aga cct act	2064
405	Ala Gln Ile Ile Leu Lys Leu Cys Phe Ser Ala Lys Phe Arg Pro Thr	
406	645 650 655	
407	aag caa aac aat caa gtg gcc att gta tat aac atc aca ctt gat gca	2112
408	Lys Gln Asn Asn Gln Val Ala Ile Val Tyr Asn Ile Thr Leu Asp Ala	
409	660 665 670 675	
410	gat gga ttt tca tcc aga gta acc tcc agg ggg tta ttt aaa gaa aac	2160
411	Asp Gly Phe Ser Ser Arg Val Thr Ser Arg Gly Leu Phe Lys Glu Asn	
412	680 685 690	
413	aat gaa agg tgc ctg cag aag aat atg gta gta aat caa gca cag agt	2208
414	Asn Glu Arg Cys Leu Gln Lys Asn Met Val Val Asn Gln Ala Gln Ser	
415	695 700 705	
416	tgc ccc gag cac atc att tat ata cag gag ccc tct gat gtt gtc aac	2256
417	Cys Pro Glu His Ile Ile Tyr Ile Gln Glu Pro Ser Asp Val Val Asn	
418	710 715 720	
419	tct ttg gat ttg cgt gtg gac atc agt ctg gaa aac cct ggc act agc	2304
420	Ser Leu Asp Leu Arg Val Asp Ile Ser Leu Glu Asn Pro Gly Thr Ser	
421	725 730 735	
422	cct gcc ctt gaa gcc tat tct gag act gcc aag gtc ttc agt att cct	2352
423	Pro Ala Leu Glu Ala Tyr Ser Glu Thr Ala Lys Val Phe Ser Ile Pro	
424	740 745 750 755	
425	ttc cac aaa gac tgt ggt gag gat gga ctt tgc att tct gat cta gtc	2400
426	Phe His Lys Asp Cys Gly Glu Asp Gly Leu Cys Ile Ser Asp Leu Val	
427	760 765 770	
428	cta gat gtc cga caa ata cca gct gct caa gaa caa ccc ttt att gtc	2448
429	Leu Asp Val Arg Gln Ile Pro Ala Ala Gln Glu Gln Pro Phe Ile Val	
430	775 780 785	
431	agc aac caa aac aaa agg tta aca ttt tca gta aca ctg aaa aat aaa	2496
432	Ser Asn Gln Asn Lys Arg Leu Thr Phe Ser Val Thr Leu Lys Asn Lys	
433	790 795 800	
434	agg gaa agt gca tac aac act gga att gtt gtt gat ttt tca gaa aac	2544
435	Arg Glu Ser Ala Tyr Asn Thr Gly Ile Val Val Asp Phe Ser Glu Asn	
436	805 810 815	
437	ttg ttt ttt gca tca ttc tcc cta ccg gtt gat ggg aca gaa gta aca	2592
438	Leu Phe Phe Ala Ser Phe Ser Leu Pro Val Asp Gly Thr Glu Val Thr	

PAGE: 10

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/155,514DATE: 02/03/2000
TIME: 13:10:50

Input Set: I155514.RAW

439	820	825	830	835
440	tgc cag gtg gct gca tct cag aag tct gtt gcc tgc gat gta ggc tac			2640
441	Cys Gln Val Ala Ala Ser Gln Lys Ser Val Ala Cys Asp Val Gly Tyr			
442	840	845	850	
443	cct gct tta aag aga gaa caa cag gtg act ttt act att aac ttt gac			2688
444	Pro Ala Leu Lys Arg Glu Gln Gln Val Thr Phe Thr Ile Asn Phe Asp			
445	855	860	865	
446	ttc aat ctt caa aac ctt cag aat cag gcg tct ctc agt ttc caa gcc			2736
447	Phe Asn Leu Gln Asn Leu Gln Asn Gln Ala Ser Leu Ser Phe Gln Ala			
448	870	875	880	
449	tta agt gaa agc caa gaa aac aag gct gat aat ttg gtc aac ctc			2784
450	Leu Ser Glu Ser Gln Glu Glu Asn Lys Ala Asp Asn Leu Val Asn Leu			
451	885	890	895	
452	aaa att cct ctc ctg tat gat gct gaa att cac tta aca aga tct acc			2832
453	Lys Ile Pro Leu Leu Tyr Asp Ala Glu Ile His Leu Thr Arg Ser Thr			
454	900	905	910	915
455	aac ata aat ttt tat gaa atc tct tcg gat ggg aat gtt cct tca atc			2880
456	Asn Ile Asn Phe Tyr Glu Ile Ser Ser Asp Gly Asn Val Pro Ser Ile			
457	920	925	930	
458	gtg cac agt ttt gaa gat gtt ggt cca aaa ttc atc ttc tcc ctg aag			2928
459	Val His Ser Phe Glu Asp Val Gly Pro Lys Phe Ile Phe Ser Leu Lys			
460	935	940	945	
461	gta aca aca gga agt gtt cca gta agc atg gca act gta atc atc cac			2976
462	Val Thr Thr Gly Ser Val Pro Val Ser Met Ala Thr Val Ile Ile His			
463	950	955	960	
464	atc cct cag tat acc aaa gaa aag aac cca ctg atg tac cta act ggg			3024
465	Ile Pro Gln Tyr Thr Lys Glu Lys Asn Pro Leu Met Tyr Leu Thr Gly			
466	965	970	975	
467	gtg caa aca gac aag gct ggt gac atc agt tgt aat gca gat atc aat			3072
468	Val Gln Thr Asp Lys Ala Gly Asp Ile Ser Cys Asn Ala Asp Ile Asn			
469	980	985	990	995
470	cca ctg aaa ata gga caa aca tct tct tct gta tct ttc aaa agt gaa			3120
471	Pro Leu Lys Ile Gly Gln Thr Ser Ser Val Ser Phe Lys Ser Glu			
472	1000	1005	1010	
473	aat ttc agg cac acc aaa gaa ttg aac tgc aga act gct tcc tgt agt			3168
474	Asn Phe Arg His Thr Lys Glu Leu Asn Cys Arg Thr Ala Ser Cys Ser			
475	1015	1020	1025	
476	aat gtt acc tgc tgg ttg aaa gac gtt cac atg aaa gga gaa tac ttt			3216
477	Asn Val Thr Cys Trp Leu Lys Asp Val His Met Lys Gly Glu Tyr Phe			
478	1030	1035	1040	
479	gtt aat gtg act acc aga att tgg aac ggg act ttc gca tca tca acg			3264
480	Val Asn Val Thr Thr Arg Ile Trp Asn Gly Thr Phe Ala Ser Ser Thr			
481	1045	1050	1055	
482	ttc cag aca gta cag cta acg gca gct gca gaa atc aac acc tat aac			3312
483	Phe Gln Thr Val Gln Leu Thr Ala Ala Ala Glu Ile Asn Thr Tyr Asn			
484	1060	1065	1070	1075
485	cct gag ata tat gtg att gaa gat aac act gtt acg att ccc ctg atg			3360
486	Pro Glu Ile Tyr Val Ile Glu Asp Asn Thr Val Thr Ile Pro Leu Met			
487	1080	1085	1090	
488	ata atg aaa cct gat gag aaa gcc gaa gta cca aca gat ccc gag			3405

PAGE: 11

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/155,514DATE: 02/03/2000
TIME: 13:10:50

Input Set: I155514.RAW

489 Ile Met Lys Pro Asp Glu Lys Ala Glu Val Pro Thr Asp Pro Glu
 490 1095 1100 1105
 491 ctgcttggaaag caggctcaggc gctccctgcct ggacgcattcc cggctatgca gccccagtc 3465
 492 agggcagcaa ggcaggcccc gtctgcctct tcacccggag cctctgccc ccccactcat 3525
 493 gctcaggagg agggctttct ggcttttcc caggctctgg gcaggcacag gctaggtgcc 3585
 494 cctaaccagg gccctgcaca caaaggggca ggtgctggc tcagacctgc caagagccat 3645
 495 atccgggagg accctgcccc tgacctaagg ccaccccaaa ggccaaactc tccactccct 3705
 496 cagctcggac accttctctc ctccagatt ccagtaactc ccaatcttct ctctgca 3762
 497 gag ccc aaa tct tgt gac aaa act cac aca tgc cca ccg tgc cca 3807
 498 Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro
 499 1110 1115 1120
 500 ggtaaggccag cccaggccctc gcccctccagg tcaaggcggg acaggtgccc tagagtagcc 3867
 501 tgcattccagg gacaggcccc agccgggtgc tgacacgtcc acctccatct cttcctca 3925
 502 gca cct gaa ctc ctg ggg gga ccg tca gtc ttc ctc ttc ccc cca aaa 3973
 503 Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys
 504 1125 1130 1135
 505 ccc aag gac acc ctc atg atc tcc cgg acc cct gag gtc aca tgc gtg 4021
 506 Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val
 507 1140 1145 1150
 508 gtg gtg gac gtg agc cac gaa gac cct gag gtc aag ttc aac tgg tac 4069
 509 Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr
 510 1155 1160 1165
 511 gtg gac ggc gtg gag gtg cat aat gcc aag aca aag ccg cgg gag gag 4117
 512 Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu
 513 1170 1175 1180 1185
 514 cag tac aac agc acg tac cgg gtg gtc agc gtc ctc acc gtc ctg cac 4165
 515 Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His
 516 1190 1195 1200
 517 cag gac tgg ctg aat ggc aag gag tac aag tgc aag gtc tcc aac aaa 4213
 518 Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys
 519 1205 1210 1215
 520 gcc ctc cca gcc ccc atc gag aaa acc atc tcc aaa gcc aaa 4255
 521 Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys
 522 1220 1225 1230
 523 ggtgggaccc gtggggtgcg agggccacat ggacagagggc cggctcgcc caccctctgc 4315
 524 cctgagagtg accgctgtac caacctctgt cctaca ggg cag ccc cga gaa cca 4369
 525 Gly Gln Pro Arg Glu Pro
 526 1235
 527 cag gtg tac acc ctg ccc cca tcc cgg gat gag ctg acc aag aac cag 4417
 528 Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln
 529 1240 1245 1250
 530 gtc agc ctg acc tgc ctg gtc aaa ggc ttc tat ccc agc gac atc gcc 4465
 531 Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala
 532 1255 1260 1265
 533 gtg gag tgg gag agc aat ggg cag ccg gag aac aac tac aag acc acg 4513
 534 Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr
 535 1270 1275 1280 1285
 536 cct ccc gtg ctg gat tcc gac ggc tcc ttc ctc tac agc aag ctc 4561
 537 Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu
 538 1290 1295 1300

PAGE: 12

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/155,514DATE: 02/03/2000
TIME: 13:10:50

Input Set: I155514.RAW

539	acc gtg gac aag agc agg tgg cag cag ggg aac gtc ttc tca tgc tcc	4609
540	Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser	
541	1305 1310 1315	
542	gtg atg cat gag gct ctg cac aac cac tac acg cag aag agc ctc tcc	4657
543	Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser	
544	1320 1325 1330	
545	ctg tct ccg ggt aaa tga	4675
546	Leu Ser Pro Gly Lys	
547	1335	

PAGE: 13

VERIFICATION SUMMARY
PATENT APPLICATION US/09/155,514

DATE: 02/03/2000

TIME: 13:10:50

Input Set: I155514.RAW

Line ? Error/Warning

Original Text

180 E Number of Bases conflict w/ Running Total atg gta cca aat tct ttt agc ccc caa act g
284 E Number of Bases conflict w/ Running Total ggt ctc cca gaa gca aaa ata ttt tcc ggt c